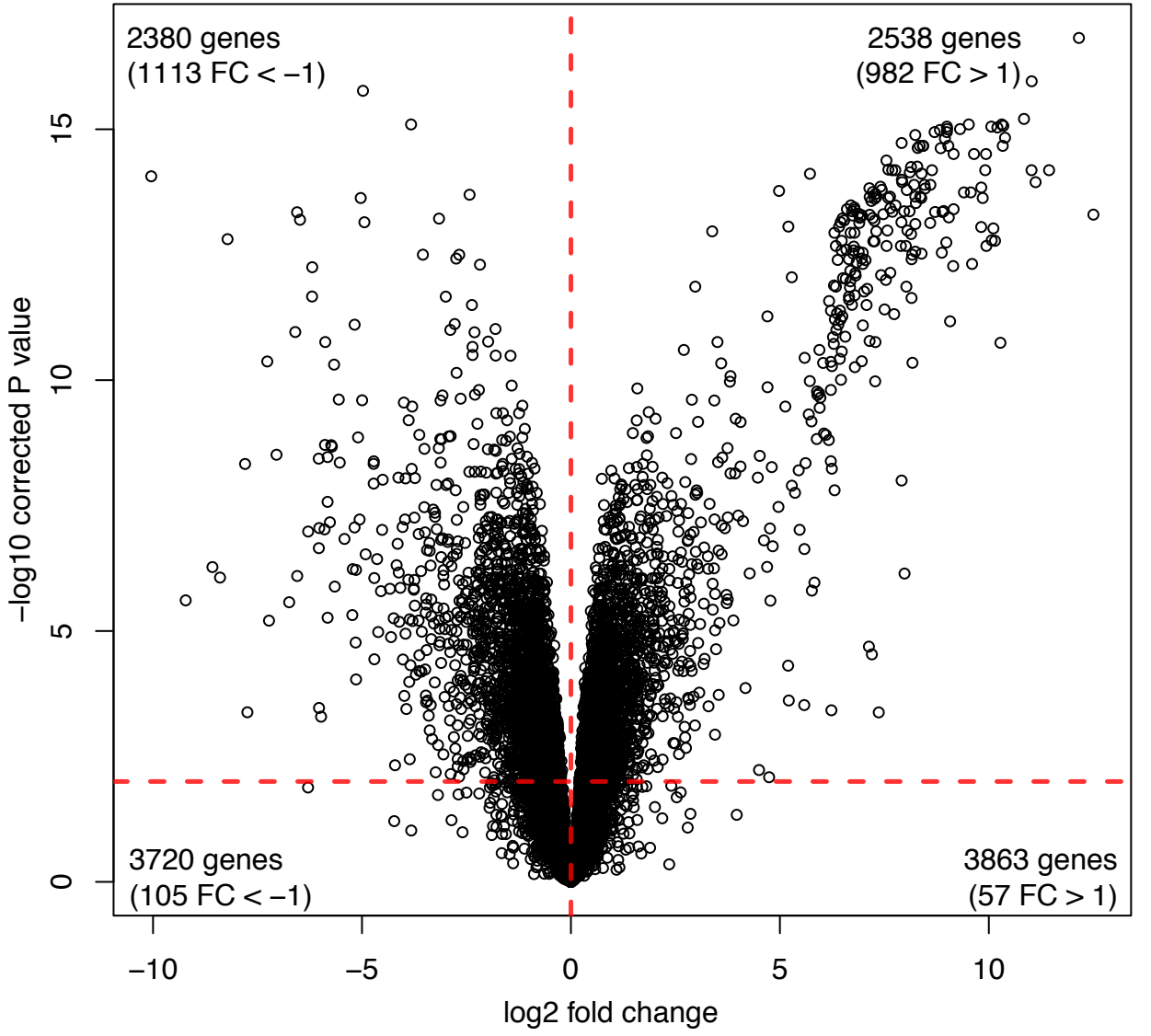
2380 genes
(1113 FC < −1)
2538 genes
(982 FC > 1)
3720 genes
(105 FC < −1)
3863 genes
(57 FC > 1)
−log10 corrected P value
0
5
10
15
−10
−5
0
5
10
log2 fold change